

## SEQUENCE LISTING

<110> Kramer, Michael

<120> Regulatory Protein pKe#83 from Human  
Keratinocytes

<130> km-3/PCT

<140> PCT/DE99/03732

<141> 1999-11-19

<150> DE19854672.6

<151> 1998-11-26

<150> DE19856301.9

<151> 1998-12-07

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<170> PatentIn Ver. 2.1

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 Caseine kinase, 2x Tyrosine kinase

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 Arg Ser Leu Glu Cys Arg Ser Asp Pro Glu Ser Pro Ile Lys Lys Thr

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Pro Asp Ala Asp Arg Thr Thr Leu Asn His Ala Asp His Ser Ser Lys		
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Pro Ser Ser Ala Ala Gln Lys Ala Val Thr Glu Ser Ser Glu Gln Asp		
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Met Lys Ser Gly Thr Glu Asp Leu Arg Thr Glu Arg Leu Gln Lys Thr		
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<223> Phoshorylation sites: 9x Protein kinase, 15x  
Caseine kinase, 2x Tyrosine kinase

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&lt;223&gt; Prenylation site (CAAX-box)

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260	265	270
Tyr Ile Glu Asn Arg Pro Glu Met Lys Arg Gln Arg Ser Ile Gln Glu		
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 Asp Leu Ala Lys Lys His Ala Ser Leu Arg Gln Thr Glu Ser Asp  
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 Pro Asp Ala Asp Arg Thr Thr Leu Asn His Ala Asp His Ser Ser Lys  
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 Ile Val Gln His Arg Leu Leu Ser Arg Gln Glu Leu Lys Glu Arg  
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Ala Arg Val Leu Leu Glu Gln Ala Arg Arg Asp Ala Ala Leu Lys Ala  
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Leu Ser Asp Gln Gln Asp Glu Glu Arg Arg Arg Gln Leu Arg Glu Arg  
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Ala Arg Gln Leu Ile Ala Glu Ala Arg Ser Gly Val Lys Met Ser Glu  
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Leu Pro Ser Tyr Gly Glu Met Ala Ala Glu Lys Leu Lys Glu Arg Ser  
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Lys Ala Ser Gly Glu Gln Asn Ser Lys Leu Val Asp Leu Lys Leu Lys  
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Lys Leu Leu Glu Val Gln Pro Gln Val Ala Asn Ser Pro Ser Ser Ala  
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Ala Gln Lys Ala Val Thr Glu Ser Ser Glu Gln Asp Met Lys Ser Gly  
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Thr Glu Asp Leu Arg Thr Glu Arg Leu Gln Lys Thr Thr Glu Arg Phe  
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Arg Asn Pro Val Val Phe Ser Lys Asp Ser Thr Val Arg Lys Thr Gln  
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Leu Gln Ser Phe Ser Gln Tyr Ile Glu Asn Arg Pro Glu Met Lys Arg  
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Gln Arg Ser Ile Gln Glu Asp Thr Lys Lys Gly Asn Glu Glu Lys Ala  
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Met Gln Ser Leu Ala Ser Leu Val Ser Met Lys Gln Ala Asp Ile Gly  
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Asn Leu Asp Asp Phe Glu Glu Asp Asn Glu Asp Asp Asp Glu Asn Arg  
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Val Asn Gln Glu Glu Lys Ala Ala Lys Ile Thr Glu Leu Ile Asn Lys  
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Leu Asn Phe Leu Asp Glu Ala Glu Lys Asp Leu Ala Thr Val Asn Ser  
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Asn Pro Phe Asp Asp Pro Asp Ala Ala Glu Leu Asn Pro Phe Gly Asp  
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Pro Asp Ser Glu Glu Pro Ile Thr Glu Thr Ala Ser Pro Arg Lys Thr  
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Glu Asp Ser Phe Tyr Asn Asn Ser Tyr Asn Pro Phe Lys Glu Val Gln  
145 150 155 160

Thr Pro Gln Tyr Leu Asn Pro Phe Asp Glu Pro Glu Ala Phe Val Thr  
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Ile Lys Asp Ser Pro Pro Gln Ser Thr Lys Arg Lys Asn Ile Arg Pro  
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Val Asp Met Ser Lys Tyr Leu Tyr Ala Asp Ser Ser Lys Thr Glu Glu  
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Glu Glu Leu Asp Glu Ser Asn Pro Phe Tyr Glu Pro Lys Ser Thr Pro  
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Pro Pro Asn Asn Leu Val Asn Pro Val Gln Glu Leu Glu Thr Glu Arg  
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Arg Val Lys Arg Lys Ala Pro Ala Pro Pro Val Leu Ser Pro Lys Thr  
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Gly Val Leu Asn Glu Asn Thr Val Ser Ala Gly Lys Asp Leu Ser Thr  
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Ser Pro Lys Pro Ser Pro Ile Pro Ser Pro Val Leu Gly Arg Lys Pro  
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Tyr Asp Gly Phe Ala Ser Ile Gly Ile Ser Arg Leu Leu Glu Pro Ser  
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Asp Met Val Leu Leu Ala Ile Pro Asp Lys Leu Thr Val Met Thr Tyr  
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Leu Tyr Val Ser Asp Lys Lys Asp Met Ser Pro Pro Phe Ile Cys  
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Val Gly Gly Gly Asp Glu Leu Thr Asn Leu Glu Asn Asp Leu Asp Thr			
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Cys Val Leu Gln  
1075

1

26